From: Graser, Jennifer AU 1645
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Ngo, J.T. et al., "Computational Complexity, Protein Structure Prediction, and the Levinthal Paradox", The Protein Folding Problem and Tertiary Structure Prediction, 1994, Merz et al (eds), Birkhauser, Boston, MA, pp. 433 and 492-495.

Mikayama et al, Molecular Cloning and Functional Expression of a cDNA encoding glycosylation-inhibiting factor, November 1993, Proc. Nat. Acad. Sci., USA, vol 90: 10056-10060.
"human glycosylation-inhibiting factor (GIF) protein differs from human macrophage migration inhibitory factor (MIF) by a single amino acid residue (Figure 1) and GIF is unable to carry out the function of MIF and MIF does not demonstrate GIF bioactivity (ABST)."